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STATISTICAL ANALYSIS PLAN

A Phase 1b Open Label Study Investigating the Safety and Efficacy of Blinatumomab in Combination With Pembrolizumab in Adult Subjects With Relapsed or Refractory Diffuse Large B Cell Lymphoma (DLBCL)

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Authors: , Global Biostatistics Science

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Table of Abbreviations

Abbreviation/Acronym	Definition		
ADPC	Analysis Dataset for PK Concentrations		
AE	Adverse Event		
AMQs	Amgen MedDRA Queries		
CDM	Clinical Data Management		
CR	Complete Response		
CPMS	Clinical Pharmacology Modeling and Simulation		
CRF	Case Report Form		
CSR	Clinical Study Report		
CTCAE	Common Terminology Criteria for Adverse Events		
DLBCL	Diffuse Large B-Cell Lymphoma		
DLRT	Dose Level Review Team		
DLT			
	Dose Limiting Toxicity		
DOR	Duration of Response		
DRE	Disease Related Events		
ECG	Electrocardiogram		
ECI	Event of Clinical Interest		
EOI	Events of interest		
E-R analysis	Exposure Response Analysis		
FAS	Full Analysis Set		
GSO-DM	Global Study Operations-Data Management		
HSCT	Hematopoietic Stem Cell Transplantation		
IA	Interim Analysis		
IP	Investigational Product		
IPD	Important Protocol Deviations		
IPI	International Prognostic Index		
IVRS	Interactive Voice Response System		
KM	Kaplan-Meier		
MedDRA	Medical Dictionary for Regulatory Activities		
MTD	Maximum Tolerated Dose		
ORR	Objective Response Rate		
OS	Overall Survival		
PFS	Progression Free Survival		
PK	Pharmacokinetics		



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Abbreviation/Acronym	Definition
PR	Partial Response
PRO	Patient Reported Outcome
SAP	Statistical Analysis Plan
SMQs	Standard MedDRA Queries
SSAP	Supplemental Statistical Analysis Plan
WHODRUG	World Health Organization Drug



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1. Introduction

The purpose of this Statistical Analysis Plan (SAP) is to provide details of the statistical analyses that have been outlined within the protocol for the phase 1b blinatumomab in combination with pembrolizumab study 20150290, dated August 21 2017. The scope of this plan includes the interim analyses, primary analysis and the final analysis that are planned and will be executed by the Global Biostatistics Science department unless otherwise specified. Pharmacokinetic, pharmacodynamic, exposure-response and biomarker analyses will be performed by Clinical Pharmacology Modeling and Simulation (CPMS) or biomarker group.

2. Objectives

2.1 Primary

To determine the maximum tolerated dose (MTD) of blinatumomab in combination with pembrolizumab in adult subjects with relapsed or refractory (r/r) DLBCL

2.2 Secondary

To evaluate the safety, efficacy, and pharmacokinetics (PK) of blinatumomab in combination with pembrolizumab in adult subjects with r/r DLBCL

2.3 Exploratory

- To evaluate blood and tissue biomarkers
- To evaluate minimal residual disease (MRD) response by next generation sequencing (NGS)
- To estimate the impact of pembrolizumab in combination with blinatumomab on patient reported outcomes (PROs)

3. Study Overview

3.1 Study Design

This is an open label, multicenter, phase 1b study testing the combination of blinatumomab with pembrolizumab in r/r DLBCL.

The study will consist of 2 portions:

- Part 1 (n = 6 50) will test the safety of up to 3 different blinatumomab target dose levels and up to 3 schedules of blinatumomab in combination with pembrolizumab in a rolling 6 design. A Dose Level Review Team (DLRT) will review the safety data to evaluate possible drug effects and dose limiting toxicities (DLTs). Subjects who are not on the dose ultimately selected for part 2 will remain on their initial dose throughout the study.
- Part 2 (n = 36) will consist of an expansion cohort to assess PK, safety, and preliminary efficacy data at the chosen target dose. The part 2 dose will be determined by the totality of the clinical data from part 1 as determined by the DLRT.



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The study design includes:

A 21-day screening period.

- A standard (core) treatment period of blinatumomab (first cycle) of 8 weeks
- A second (consolidation) cycle of blinatumomab of 28 days after a 28-day (± 3 days) blinatumomab treatment-free period that can be administered to subjects with stable disease (SD), partial response (PR), or complete response (CR).
- Pembrolizumab treatment until disease progression or up to 35 cycles in the absence of disease progression:
 - On study day 15 for subjects in cohort la

OR

On study day 1 for subjects in cohorts lb, llb, and lllb

OR

- On study day 19 for subjects in cohort IIa and IIIa
- A safety follow-up visit after 30 days (+ 7 days) of last dose of each protocol specified therapy.

Follow-up for survival and collection of subsequent anticancer therapies will occur every 12 weeks (± 28 days) following blinatumomab safety follow-up visit for up to approximately 24 months from the last dose of pembrolizumab.

Part 1 design and blinatumomab escalation/de-escalation rules

For part 1, subject enrollment will start in cohort la as outlined in the schema in Figure 1. Blinatumomab will be dosed as a continuous intravenous infusion (CIVI) for 8 weeks. The initial dose will be 9 μ g/day and the dose will be escalated after 7 days to a target dose of 28 μ g/day. Depending on tolerability, the target dose of blinatumomab will be increased to a maximum of 112 μ g/day in cohort IIa and IIb, with possible de-escalation to 56 μ g/day in cohorts IIIa and IIIb. Pembrolizumab will be dosed by intravenous (IV) infusion 200 mg at Q3W starting on study day 15 in cohort Ia, starting on study day 1 in cohorts Ib, IIb, and IIIb, and on study day 19 in cohorts IIa and IIIa.

Subjects who do not meet the criteria for investigational product (IP) discontinuation (see below) are eligible for a second cycle of blinatumomab (consolidation) consisting of a CIVI of 28 days after a 28-day (± 3 days) blinatumomab treatment-free interval. Blinatumomab will be started at 9 µg/day and escalated every 7 days to the maximum target dose of blinatumomab in the assigned cohort.

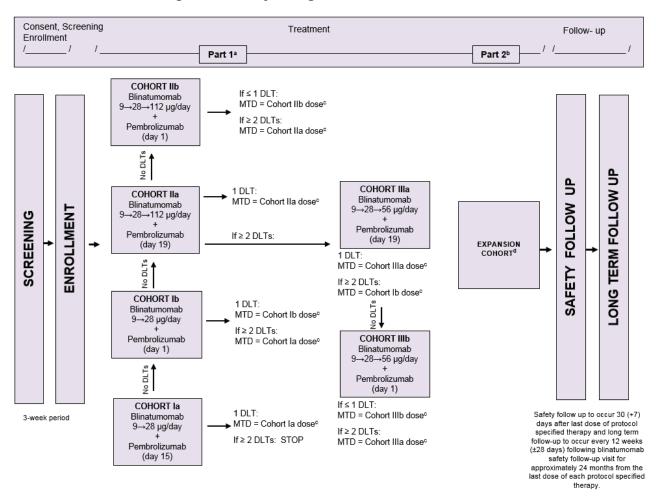


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Study Design and Treatment Schema

Figure 1. Study Design and Treatment Schema





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DLT = dose limiting toxicity; MTD = maximum tolerated dose

First cycle of blinatumomab will be 8 weeks in duration, followed by a 28-day (± 3 days) blinatumomab treatment-free interval. A second consolidation cycle of blinatumomab will be 28 days in duration at the same dose as the first cycle, starting at 9 µg/day with weekly dose escalations until the target dose is reached, if subject has stable disease or partial/complete response after cycle 1. Pembrolizumab will be started on study day 15 for cohort Ia, study day 1 for cohorts Ib, Ilb, and IIIb, and study day 19 for cohorts IIa and IIIa, and administered Q3 weeks until disease progression for up to 35 cycles.

- ^a Part 1: To determine maximum tolerated dose (MTD) of blinatumomab in combination with pembrolizumab. The MTD will be defined as the dose level at which ≤ 1 of 6 subjects experience a Dose limiting toxicity (DLT) or the maximum administered dose (MAD).
- ^b Part 2: Expansion cohort to estimate the efficacy of the combination of blinatumomab and pembrolizumab. Dosing will be determined based on the MTD of blinatumomab established in part 1. DLTs will be continuously monitored to ensure they do not reach a pre-defined threshold.
- ^c For cohorts Ia, IIa and IIIa, the DLT observation period will begin on the same day as the first dose of pembrolizumab (day 15 for Ia and day 19 for IIa and IIIa) and will continue for 42 days. For cohort Ib, the DLT observation period will begin on day 1 of the start of the combination of pembrolizumab/blinatumomab, and continue for 42 days. For cohorts IIb, and IIIb, the DLT observation period will begin once the blinatumomab target dose (28 μg/day on day 8, 112 μg/day on day 15, or 56 μg/day on day 15 for cohorts Ib, IIb, and IIIb, respectively) is reached and will continue for 28 days. A dose level review team (DLRT) will review the available data to determine if blinatumomab is safe and tolerable as defined by DLT criteria.
- ^d Dosing for the Part 2 expansion cohort will be based on the safety of the combination of blinatumomab and pembrolizumab and the MTD of blinatumomab in Part 1.



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Subjects will be enrolled to part 1 with up to 6 subjects being enrolled per cohort. In any cohort, assuming adequate tolerability (≤ 1 DLT), up to 10 subjects may be enrolled to ensure adequate safety and PK data is collected. The decision to expand a cohort will be made by the DLRT.

The MTD of blinatumomab will be defined as the dose level at which at most 1 of 6 subjects experiences a DLT or the maximum administered dose (MAD). The MAD to be tested will be 112 µg/day (cohort IIa and IIb). The MTD defines the stopping rules for the study. Subjects who discontinue treatment prior to reaching the target dose in part 1 will be replaced. The DLT criteria is defined in the Appendix E of the protocol.

Part 2

For part 2, the dosing will be determined based on the safety of the combination of blinatumomab and pembrolizumab and the MTD of blinatumomab established in part 1 per DLRT. Part 2 will consist of an expansion cohort to collect further safety and PK data as well as provide a preliminary estimate of the efficacy of the combination of blinatumomab and pembrolizumab. Dose limiting toxicities will be monitored to ensure they do not reach a pre-defined threshold of 25%. If this threshold is reached, the DLRT will have the discretion to change to another dose/schedule tested in phase 1 part 1 based on the totality of the available data. The details of DLT boundaries are provided in Section 8.1.

3.2 Sample Size

Part 1:

A rolling 6 dose design will be used. An additional 4 subjects per cohort can be enrolled to further evaluate safety and PK data if needed. There will be a minimum of 6 subjects and a maximum of 50 subjects enrolled.

Part 2:

The sample size for part 2 is determined by a 1-sample test of ORR within 12 weeks after starting blinatumomab. With the 1-sided type I error rate (α) set at 0.05, a null hypothesis response probability (π 0) of 15%, and an alternative response probability $(\pi 1)$ of 35%, a sample size of 42 subjects (including 6 subjects enrolled in part 1 at the chosen dose level) will provide 90% power to reject the null hypothesis that the response probability is no more than 15%.



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The following parameters are used:

- H0: $\pi \le \pi 0 = 15\%$ vs H1: $\pi \ge \pi 1 = 35\%$.
- Overall type I error (alpha) for the one-sided hypothesis = 5%
- Power = 90%

If 11 of the 42 subjects achieve a response, then the null hypothesis that ORR is less than or equal to 15% can be rejected.

Study Endpoints and Covariates 4.

4.1 **Study Endpoints**

4.1.1 **Primary Endpoints**

o Incidence of DLTs

4.1.2 **Secondary Endpoints**

- o ORR (including CR and PR) within 12 weeks after starting blinatumomab by Cheson (2007) Criteria
- CR rate within 12 weeks after starting blinatumomab by Cheson (2007) Criteria
- o PFS
- OS
- Duration of response (DOR) by ORR, CR, and PR within 12 weeks after starting blinatumomab
- Blinatumomab PK parameters
- Pembrolizumab PK parameters

4.1.3 **Safety Endpoints**

Incidence and severity of adverse events

4.1.4 **Exploratory Endpoints**

- ORR by Lugano (2014) Classification
- PD-L1 expression on tumor
- Changes in Lymphocytes (B-cell, T-cell populations, NK cells) and leukocyte populations (leukocytes, lymphocytes, monocytes, and granulocytes) in peripheral blood
- Peripheral blood cytokine levels
- MRD by NGS after cycle 1 of blinatumomab
- Changes in PROs (FACT-Lymphoma, EQ-5D) from baseline



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5. Hypotheses and/or Estimations

Part 1:

The maximum tolerated dose (MTD) of blinatumomab in combination with pembrolizumab will be determined in adult subjects with r/r DLBCL.

Part 2:

Statistical hypothesis for ORR by Cheson (2007) within 12 weeks after starting blinatumomab

H0: $\pi \le \pi_0 = 15\%$ vs H₁: $\pi \ge \pi_1 = 35\%$.

6. Definitions

6.1 General Definitions

Age at Enrollment

Subject age at enrollment will be collected in years in the clinical database.

Baseline

For the analysis of all endpoints, baseline will be defined as the value measured on day 1 of the first cycle of blinatumomab. The protocol specifies that procedures and labs on day 1 should be completed before the initiation of any protocol-specified therapy which will be the assumption in the analysis unless the time of the assessment is recorded. If a day 1 value is not available, the most recent value before the day of the start of any protocol-specified therapy may be used.

Cumulative Dose of Blinatumomab, Pembrolizumab

Blinatumomab: The cumulative dose in µg is defined as the following with summation over infusions:

Σ (duration of infusion (days) for each dose received × dose received [μg])

Pembrolizumab: The cumulative dose in mg is defined as the following with summation over infusions:

(duration of Infusion (days) for each dose received × dose Received [mg])

Cumulative dose will be calculated within a cycle and across all cycles.



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Death Date

For subjects who die during the study, the death date will be recorded on the event, end of study and survival status CRF. The earliest date will be used if the dates are inconsistent among these CRF pages. For deaths collected after a subject has ended study (eq, through public records in countries where permitted), the death date will be recorded on the long term follow-up page.

Dose limiting toxicity (DLT)

The occurrence of any of the toxicities in the Appendix E of the protocol will be considered a DLT, if judged by the investigator to be possibly, probably or definitely related to study drug administration.

DLT Evaluable

To be DLT evaluable, subjects must meet one of the following criteria:

The subject experienced a DLT in the DLT evaluation period;

OR

The subject was removed from treatment for an adverse event/toxicity;

OR

The subject was removed from treatment for reasons other than an adverse event/toxicity (ie, disease progression), and the subject has received at least 12 days of pembrolizumab treatment with blinatumomab at the target dose;

OR

The subject did not experience a DLT and completed the DLT observation period.

Duration of Blinatumomab/Pembrolizumab

Blinatumomab/ Pembrolizumab: For each infusion episode within a cycle, the duration of exposure will be calculated by subtracting the start date and time from the stop date and time. If either a start or stop time is missing, only the date portion will be used in calculating the duration of a specific infusion. For each cycle, the duration will be last date minus first date plus 1 of infusion. For the entire study, the duration will be the sum of the durations across cycles. The duration will be rounded to the nearest day.



Event of Clinical Interest (ECI)

Selected non-serious and serious adverse events are known as ECI and must be reported within 24 hour to the sponsor. The ECI for this trial includes

- An overdose of blinatumomab or pembrolizumab (see Protocol Section 6.2.1.1 and Section 6.2.2.1)
- Hepatic disorder(See protocol section 9.1.4)

Events of interest (EOI)

Events of interest (EOI) will be based on search strategies defined by Standard MedDRA Queries (SMQs) or Amgen MedDRA Queries (AMQs).

End of Investigational Product (IP) Administration Date

End of IP Admininstration for subjects who had blinatumomab or pembrolizumab is defined as the last infusion of blinatumomab/ pembrolizumab reported on the End of IP Administration CRF.

Enrollment Date

Enrollment Date is defined as the date of enrollment collected on the CRF.

Investigational Product

Amgen Investigational Product: refers to blinatumomab

Non-Amgen Investigational Product: refers to pembrolizumab

<u>Last Dose Date of Blinatumomab/ Pembrolizumab</u>

This is the stop date of the last infusion of blinatumomab/pembrolizumab administration reported on the Investigational Product Administration CRF.

Percent of Intended Dose of Blinatumomab

For a cycle, the percent of intended dose of blinatumomab will be the cumulative dose, in micrograms, in that cycle divided by the planned cumulative dose for that cycle.

For the entire study, the percent of intended dose of blinatumomab will be the sum of the cumulative doses across cycles divided by the sum of the planned cumulative doses across the cycles started. Re-started cycles will have the planned cumulative dose counted both for the period before the re-start and for the period after the re-start in the calculation of the percent of intended dose.



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Prior Salvage Regimens

Prior salvage regimens are those medications recorded on the prior anti-cancer therapies CRF where the line of therapy field indicates 2nd or higher line thereapy.

Study Day

Study Day 1 is defined as the day of first dose of blinatumomab.

And Study Day is defined as:

Pre study day 1: study day= (date – date of study day 1)

Post study day 1: study day= (date - date of study day 1) + 1

Subject Level End of Study (EOS) Date

End of Study for each subject is defined as the date the subject last completed a protocol-specified procedure. The date will be recorded on the End of Study CRF page.

Treatment Emergent Adverse Event (AE)

Treatment emergent adverse event refers to an adverse event that starts on or after first dose of blinatumomab up to and including 30 days after the end of blinatumomab or pembrolizumab whichever is later. It is indicated by the flag whether an event start before first dose of blinatumomab on the Event CRF page. This reporting window also applies to treatment-emergent serious adverse events (SAEs).

<u>Treatment Emergent Disease-related Event (DRE)</u>

Treatment emergent disease-related event refers to a disease-related event that starts on or after first dose of blinatumomab up to and including 30 days after the end of blinatumomab or pembrolizumab whichever is later. It is indicated by the flag whether an event start before first dose of blinatumomab on the Event CRF page.

<u>Treatment Emergent Event of Clinical Interest (ECI)</u>

ECI is defined in Section 9.1.4 of the protocol. Treatment emergent ECI refers to an ECI that starts on or after first dose of blinatumomab up to and including 30 days after the end of blinatumomab or pembrolizumab whichever is later.



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6.2 Efficacy Endpoints

Complete Response (CR) Rate

Complete response rate is the proportion of subjects who have achieved complete response within 12 weeks after starting blinatumomab for phase 1b among all subjects in the respective analysis set.

Duration of Response (DOR)

Duration of response will be calculated only for subjects who achieve an ORR, CR or PR by Cheson (2007) criteria within 12 weeks after starting blinatumomab. The duration will be calculated from the date a response, CR or PR, is first achieved until the earliest date of a disease assessment indicating a disease progression or death, whichever occurs first. Subjects who do not have a relapse event will be censored on their last disease assessment date. If the last disease assessment date is after the date that triggers the analysis, the subject will be censored at the analysis trigger date. A sensitivity analysis will censor subjects who received HSCT at the time of HSCT unless there is no assessment after the HSCT, in which case the last assessment prior to the HSCT will be used as the censoring time.

Overall Response (OR)

OR (CR or PR) is determined by central assessment by Cheson (2007) criteria. OR determined by Lugano (2014) classification is an exploratory endpoint.

Objective Response Rate (ORR)

Objective response rate (ORR) is the proportion of subjects who have achieved either a CR or a PR among subjects within 12 weeks after starting blinatumomab in the respective analysis set.

Overall Survival (OS)

The overall survival will be calculated as the time from the date of first dose of blinatumomab until death due to any cause. Subjects who are alive at the date that triggers the analysis will be censored at the date last known to be alive. If the date last known to be alive is after the date that triggers the analysis, the subject will be censored at the analysis trigger date.



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Progression Free Survival (PFS)

The PFS will be calculated as the time from the date of first dose of blinatumomab until the date of diagnosis of progression of lymphoma per central review, or date of death, whichever is earliest. Subjects who are alive and did not have progression will be censored at the last date of tumor assessment. Progression free survival for subjects who were enrolled in dose cohorts that were not selected for the phase 1b extension part will not be calculated.

6.3 **PRO Endpoint**

PRO endpoint will be specified in a separate Supplemental SAP (SSAP).

7. **Analysis Subsets**

7.1 **Full Analysis Set**

The full analysis set includes all subjects who received blinatumomab.

7.2 Safety Analysis Set

The safety analysis set includes all subjects who received blinatumomab or pembrolizumab.

7.3 Responder Analysis Set

The responder analysis set includes all subjects who had CR or PR within 12 weeks after starting blinatumomab.

7.4 **DLT Analysis Set**

DLT analysis set includes all subjects who are DLT-evaluable.

7.5 **Pharmacokinetic Analysis Set**

The pharmacokinetic analysis set for blinatumomab includes all subjects who received any infusion of blinatumomab and have at least 1 pharmacokinetic sample collected.

The pharmacokinetic analysis set for pembrolizumab includes all subjects who received any infusion of pembrolizumab and have at least 1 pharmacokinetic sample collected.

7.6 **Pharmacodynamic Analysis Set**

The cytokine analysis set includes all subjects who receive any infusion of blinatumomab and have at least 1 cytokine sample collected.

For other biomarker analysis set include all subjects who received blinatumomab and pembrolizumab and have at least one biomarker sample collected.

7.7 Interim Analyses Set(s)

Interim analysis set will include all subjects at the interim analysis.



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8. Interim Analysis and Early Stopping Guidelines

8.1 Interim Analyses

Amgen will conduct evaluations of the ongoing DLT rate to assess if the threshold for early trial termination has been reached.

The stopping rules use a Bayesian approach to terminate the study if the posterior probability that the DLT rate is greater than 25% is > 90%. The stopping boundaries assume a prior beta distribution (0.50, 1.50). The evaluations could occur more frequently if necessary to address emerging safety concerns. The operating characteristics in Table 1 provide the probability of stopping the trial early for given hypothetical true DLT rates whereas the stopping criteria in Table 2 are based on situations where the empirical evidence would result in a posterior probability of ≥90% that the true DLT rate is \geq 25%.

Table 1. Stopping Boundary With Batch Size of 7 Subjects, Posterior Probability of 90% and DLT Limit of 25%

Number of DLT evaluable subjects	Stop study if observing these many DLTs
7	≥ 4
14	≥ 6
21	≥ 9
28	≥ 11
35	≥ 13
42	Study completes

DLT = dose limiting toxicity

Table 2. Operating Characteristics With Batch Size of 7 Subjects

True DLT Rate	Prob of Stopping	Average Sample Size	
0.20	7%	40	
0.25	19% 37		
0.30	37%	33	
0.35	59%	27	
0.40	78%	22	

DLT = dose limiting toxicity; prob = probability

8.2 **Dose Level Review Team**

A DLRT will review safety data from each cohort in part 1 to recommend if blinatumomab and pembrolizumab in combination is safe and tolerable as defined by DLT criteria, taking into account a general benefit risk assessment. Pharmacokinetic data may be



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reviewed, if available. The DLRT will meet to confirm the decision rules when any of the following criteria are met:

• 2 or more subjects have experienced a DLT in a cohort

- 6 subjects are enrolled in a cohort and all subjects have completed the DLT observation period
- In the event that a cohort is expanded to 10, DLRT may also meet after all subjects have completed DLT observation period

The DLRT will consist of, at minimum, members from the Amgen study teams, including at least one clinician, one safety representative, one statistician, at least one representative of the Merck study team, and one investigator participating in the study who has recruited subjects into the cohort under review. A cohort may be expanded by DLRT recommendation in the case that the data suggests a change to the anticipated risk/benefit profile, warranting collection of further data at the blinatumomab target dose.

In part 2, DLRT will meet after every set of 7 subjects who become DLT evaluable.

9. Data Screening and Acceptance

9.1 General Principles

The objective of the data screening is to assess the quantity, quality, and statistical characteristics of the data relative to the requirements of the planned analyses. The database will be subject to edit checks outlined in the data management plan by Amgen Clinical Data Management (CDM) department. Any outstanding data issues will be communicated to CDM for resolution before the database is locked.

9.2 Data Handling and Electronic Transfer of Data

The Amgen Global Study Operations-Data Management (GSO-DM) department will provide all data to be used in the planned analyses. This study will use the RAVE Clinical database.

An Analysis Dataset for PK Concentrations (ADPC) will be provided to the appropriate Clinical Pharmacology Modeling and Simulation CPMS representative from Global Biostatistical Sciences.

9.3 Handling of Missing and Incomplete Data

Subjects without tumor response assessments will be considered as nonresponders. Otherwise, only nonmissing data will be analyzed. No other missing value replacement procedure will be deployed for clinical data. The handling of incomplete and partial dates for adverse events and concomitant medications are described in Appendix A.



Handling of missing or incomplete data for exposure-response analysis will be described in the E-R supplemental SAP (SSAP) or associated documents to support population PK/PD dataset generation and E-R analysis.

9.4 **Detection of Bias**

Methods to detect bias are described in the analyses of particular endpoints.

9.5 **Outliers**

Any suspected outliers will be investigated by the study team and will be included in the database unless determined to be an error or there is supporting evidence or explanation to justify the exclusion. Any outliers excluded from the analysis will be discussed in the Clinical Study Report (CSR), including the reasons for exclusion and the impact of their exclusion on the study.

PK Serum concentration data will be evaluated for outliers by visual inspection, and decisions to re-assay individual samples will be made in accordance with standard pharmacokinetic evaluation practice.

9.6 **Distributional Characteristics**

The statistical assumptions for analysis methods will be assessed. If the assumptions for the distributional characteristics are not met, these will be described and further analyses may be carried out using data transformations or alternative analysis methods. The use of transformations or alternative analysis methods will be justified in the final study report.

9.7 Validation of Statistical Analyses

Programs will be developed and maintained, and output will be verified in accordance with current risk-based quality control procedures.

Tables, figures, and listings will be produced with validated standard macro programs where standard macros can produce the specified outputs.

The production environment for statistical analyses consists of Amgen-supported versions of statistical analysis software; for example, the SAS System version 9.4 or later.

10. Statistical Methods of Analysis

10.1 **General Principles**

The analysis will be performed by cohorts for part 1. The analyses for part 2 will combine subjects in part 2 and the cohort in part 1 with selected dose for part 2 unless



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specified otherwise. The analyses will be performed on the Full Analysis Set unless specified otherwise.

Continuous variables will be summarized by the nonmissing sample size n, mean, standard deviation, median, first and third quartiles, minimum, and maximum. Categorical variables will be summarized by the n and percentage in each category. Time to event endpoints will be summarized with hazard ratios, Kaplan-Meier (KM) curves, KM proportions at select time points, KM quartiles (when estimable), the number of subjects with events, the number of subjects censored, and the pattern of censoring. Point estimates for efficacy endpoints will be accompanied by 2-sided 95% confidence intervals including estimates of KM quartiles (Brookmeyer and Crowley, 1982), KM proportions (Kalbfleisch and Prentice, 1980), and binomial proportions (Clopper and Pearson, 1934). Pharmacokinetics will be performed by noncompartmental analysis. Pharmacodynamic samples will be summarized by descriptive statistics.

Relationships among drug exposures and efficacy, safety, and biomarkers may be explored if the data are sufficient.

10.2 **Subject Accountability**

The number (and percent) of subjects who were screened, received blinatumomab and pembrolizumab and complete the study will be summarized. The number (and percent) of subjects who discontinue each treatment and the study and their reasons for discontinuation will be summarized.

10.3 **Important Protocol Deviations**

Categories for Important Protocol Deviations (IPDs) are defined by the study team before the first subject's visit and updated during the IPD reviews throughout the study prior to database lock. These definitions of IPD categories, sub-category codes, and descriptions will be used during the course of the study. Eligibility deviations are defined in the protocol.

10.4 **Demographic and Baseline Characteristics**

Demographic (ie, age, age group, sex, race, ethnicity) and baseline disease characteristics will be summarized using descriptive statistics. If multiple races have been reported for a subject, the subject will be categorized as multiple race as well as by combination of races.



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The baseline characteristics to be summarized include:

- Age
 - age group: <65, 65-74, ≥75
 - summary statistics
- Sex: Male, Female
- Race
 - American Indian or Alaska Native
 - Asian
 - Black (or African American)
 - Native Hawaiian or Other Pacific Islander
 - White
 - Other
- Geographic region
 - US/Canada
 - Asia
 - Europe
 - Rest of the world
- aalPI: Low (0), Low-Intermediate (1), High-Intermediate (2), High (3)
- Primary disease status: Relapsed, Refractory
- Prior HSCT: Yes, No
- Cell of Origin Determination: GCB, Non-GCB, ABC, Not done
- Bcl-2 rearrangement status: Yes, No, Not done
- Bcl-2 overexpression status: Yes, No, Not done
- Bcl-6 rearrangement: Yes, No, Not done.
- Bcl-6 overexpression: Yes, No, Not done.
- C-myc rearrangement status: Yes, No, Not done
- C-myc overexpression status: Yes, No, Not done
- Double expresser: Yes, No (Yes is defined as both C-myc and Bcl2 expression are Yes, and both C-myc and Bcl2 rearrangement are No; otherwise No)
- Double hit: Yes, No (Yes is defined by both C-myc and Bcl-6 overexpression and rearrangement are Yes; otherwise No)

10.5 **Efficacy Analyses**

The efficacy endpoints are ORR, CR, PFS, OS and DOR.

The percentage of subjects with an objective response (CR/ PR) by Cheson (2007) criteria within 12 weeks after starting blinatumomab and during the entire treatment



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period will be summarized with an exact binomial 90% confidence interval. The percentage of subjects with an objective response (CR/PR) by Lugano (2014) criteria within 12 weeks will also be summarized. Subjects missing post baseline disease assessments will be considered not to have achieved an objective response.

The KM summaries will be performed for PFS, OS, and DOR. K-M quartiles along with 2-sided 95% CIs, the number of subjects censored and the number of subjects with events will be provided. The analyses of DOR wil be performed using Responder Analysis Set.

10.6 Safety Analyses

10.6.1 **DLT Summary**

The DLT criteria is defined in the Appendix E of the protocol.

Incidence rates of DLTs and corresponding exact binomial 95% CIs will be summarized for DLT evaluable subjects.

10.6.2 Adverse Events and Disease Related Events

The Medical Dictionary for Regulatory Activities (MedDRA) version 19.1 or later will be used to code all events categorized as adverse events (AEs) or disease-related events (DREs).

Subject incidence of events will be summarized for all TEAEs, serious TEAEs, TEAEs leading to interruption or withdrawal of blinatumomab, fatal AEs, treatment emergent DREs and fatal DREs by system organ class and preferred term in descending order of frequency.

Subject incidence of ECIs will be summarized by ECI category and preferred term in descending order of frequency.

10.6.3 **Laboratory Test Results**

Plots and summary statistics over scheduled visits for actual values, changes from baseline of selected laboratory parameters below will be presented.

- 1. Calcium
- 2. Magnesium
- 3. Total bilirubin
- 4. Direct bilirubin
- Alkaline phosphatase
- 6. AST (SGOT)



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- 7. ALT (SGPT)
- 8. Amylase
- 9. Lipase
- 10. Hemoglobin
- 11. Platelets
- 12. Neutrophils
- 13. Lymphocytes
- 14. LDH
- 15. Immunoglobulins (IgG, IgA, IgM)

10.6.4 Vital Signs

The number and percentage of subjects with abnormal changes in systolic blood pressure, diastolic blood pressure and heart rate will summarized.

10.6.5 Physical Measurements

Not Applicable

10.6.6 Electrocardiogram (ECG)

Not Applicable

10.6.7 Antibody Formation

The incidence and percentage of subjects who develop anti blinatumomab antibodies (binding and if positive, neutralizing) and anti-pembrolizumab antibodies at any time will be tabulated.

10.6.8 Exposure to Investigational Product

Descriptive statistics will be produced to describe the exposure to blinatumomab and pembrolizumob. The number of cycles initiated, completed, discontinued, and re-started of blinatumomab and pembrolizumab will be summarized. In addition, the duration of therapy will be summarized by cycle and overall. The number and percent of subjects with dose modifications (eg, dose changes, dose interruptions) and reasons for modification will be summarized.

10.6.9 Exposure to Concomitant Medication

The number and percent of subjects receiving concomitant medications from study day 1 through blinatumomab or pembrolizumab safety follow-up, whichever is later will be summarized by preferred term as coded by the World Health Organization Drug (WHODRUG) dictionary. In addition, the number and proportion of subjects receiving anti-cancer therapies during long term follow-up will be summarized.



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10.6.10 Pharmacokinetic Analysis of Blinatumomab

Blinatumomab serum samples will be taken as listed in Schedule of Assessment of protocol. All data for PK analyses will be extracted from a secure folder and were mapped via the SAS_CDISC_v2 clinical connector into the Pharsight Knowledgebase Server (PKS) system version 4.0.3 (Pharsight®, St. Louis, MO).

Blinatumomab pharmacokinetic analysis will be performed using Phoenix WinNonlin v.6.4 software on Citrix (Pharsight®, St. Louis, MO) as part of the validated PKS system on individual serum blinatumomab concentrations to estimate the following PK parameters:

- The steady state serum concentration (C_{ss}) of cycle 1 summarized as the observed concentrations collected during continuous IV infusion by dose levels.
- Systemic clearance (CL) calculated as CL=R₀/C_{ss}; where R₀ is the infusion rate (μg/hr) and C_{ss} is the dose normalized average C_{ss}.

Nominal times were used for presenting data in tables. Blinatumomab concentrations below the lower limit of quantification (LLOQ, 50 pg/mL) were set to zero before data analysis. All individual PK parameters and descriptive statistics are presented to 3 significant figures, except for CV%, which was reported to 1 decimal place.

PK parameters such as steady state concentration (Css) will be estimated for patients who have evaluable PK data. Summary statistics, including mean, standard deviation, CV%, median, range (Minimal, Maximal), geometric mean and CV% of geometric mean will be computed for each pharmacokinetic parameter and grouped by dose, and treatment Phase. Individual concentration-time data will be tabulated and presented in PK appendix. Mean concentration-time profiles for each cohort may be provided, if sufficient data are available.

10.6.11 Pharmacokinetic Analysis of Pembrolizumab

Pembrolizumab serum samples will be taken as listed in Schedule of Assessment of protocol. Pembrolizumab exposure parameters (eg, concentrations at the end of IV infusion and steady state trough concentrations) will be estimated for subjects who have evaluable PK data. Summary statistics, including mean, standard deviation, CV%, median, range (Minimal, Maximal), geometric mean and CV% of geometric mean will be computed for each pharmacokinetic parameter by treatment cycles. Individual concentration-time data will be tabulated and presented in PK appendix. Mean concentration-time profiles may be provided, if sufficient data are available.



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10.6.12 Pharmacodynamic Analysis

Blood samples for biomarker analysis will be taken as described in the Schedule of Assessment of protocol. The cytokine levels over time will be analyzed with summary statistics by treatment period and treatment cohort. Other analysis may be performed as appropriate.

Merck biomarker lead will define analysis related to pembrolizumab and PD-L1.

10.6.13 Exposure Response Analysis

PK data of blinatumomab may be subjected to exploratory population PK analysis with data from multiple studies. Nonlinear mixed effects modeling will be used for the analysis. Effect of covariates on exposure will be determined. These may include, age, body weight, body surface area, renal function, liver function, sex and selected baseline lab values. Other covariates may be analyzed as needed. Individual blinatumomab concentration data at time of interest may be used for the exposure response analysis.

Exposure-response relationships for selected efficacy and safety endpoints may be assessed as appropriate. The objectives and methodology of the exposure-response analysis will be provided in an E-R SSAP.

11. Changes From Protocol-specified Analyses

There are no changes to the protocol-specified analyses.



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12. Literature Citations / References

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Cui L, Hung HM, Wang SJ. Modification of Sample Size in Group Sequential Clinical Trials. *Biometrics*. 1999:853-857.

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13. **Prioritization of Analyses**

Not Applicable.

14. **Data Not Covered by This Plan**

Not Applicable.



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15. Appendices

Appendix A. Technical Detail and Supplemental Information Regarding Statistical Procedures and Programs

Handling Incomplete Dates and Missing Dates for Adverse Events and Concomitant Medications.

The following data will be imputed using the following algorithm:

- Adverse Events
- Concomitant Medications

Imputation Rules for Partial or Missing Start Dates

		Stop Date						
		Complete: yyyymmdd		Partial: <i>yyyymm</i>		Partial: <i>yyyy</i>		
Start Date		< 1 st dose	≥ 1 st dose	< 1 st dose <i>yyyymm</i>	≥ 1 st dose <i>yyyymm</i>	< 1 st dose <i>yyyy</i>	≥ 1 st dose <i>yyyy</i>	missing
Partial: yyyymm	= 1 st dose yyyymm	2	1	2	1	n/a	1	1
	≠ 1 st dose yyyymm		2		2	2	2	2
Partial: <i>yyyy</i>	= 1 st dose	3	1	3	1	n/a	1	1
	≠ 1 st dose		3		3	3	3	3
Missing		4	1	4	1	4	1	1

^{1 =} Impute the date of first dose

Note: If the start date imputation leads to a start date that is after the stop date, then do not impute the start date.

Imputation rules for partial or missing stop dates:

Initial imputation

- For partial stop date mmyyyy, impute the last of the month.
- For partial stop date yyyy, impute December 31 of the year.
- For completely missing stop date, do not impute.



^{2 =} Impute the first of the month

^{3 =} Impute January 1 of the year

^{4 =} Impute January 1 of the stop year

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• If the stop date imputation leads to a stop date that is after the death date, then impute the stop date as the death date.

• If the stop date imputation leads to a stop date that is before the start date, then there is a data error and do not impute the stop date. (ie. set the stop date as missing).

Imputation rules for partial or missing death dates:

- If death year and month are available but day is missing:
- If mmyyyy for last contact date = mmyyyy for death date, set death date to the day after the last contact date.
- If mmyyyy for last contact date < mmyyyy for death date, set death date to the first day of the death month.
- If mmyyyy for last contact date > mmyyyy for death date, data error and do not impute.

If both month and day are missing for death date or a death date is totally missing, do not impute.



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Appendix B. Reference Values/Toxicity Grades

Laboratory Values

Safety laboratory values below a distinct limit (eq. detection limit, documented as "< [limit]") will be substituted by half of the limit and values above a distinct limit (documented as "> [limit]") will be substituted by the limit itself for all analyses. A Grade (based on CTC AE version 4.0 [v4.03: June 14, 2010]) will be assigned to each laboratory result as detailed in Table 6. Depending on the toxicity definition, the same result may be assigned to two grading for deviations towards higher or lower values. In case no lower limit of normal is provided for the absolute lymphocyte, neutrophils or leukocyte counts it will not be differentiated between grade 1 and grade 0 results for these parameters. Values not meeting any of the criteria will be assigned a grade 0.

Grading of Select Laboratory Parameters

Laboratory				
Parameter [Unit]	Grade 1	Grade 2	Grade 3	Grade 4
Lymphocytes [G/L]	0.8 - < LLN	0.5 - < 0.8	0.2 - < 0.5	< 0.2
Neutrophils [G/L]	1.5 - < LLN	1.0 - < 1.5	0.5 - < 1.0	< 0.5
Leukocytes [G/L]	3.0 - < LLN	2.0 - < 3.0	1.0 - < 2.0	< 1.0
Platelets [G/L]	75 - < LLN	50 - < 75	25 - < 50	< 25
Hemoglobin [g/L]*	100 - < LLN	80 - < 100	65 - < 80	< 65
Albumin [g/L]	30 - < LLN	20 - < 30	< 20	not defined
AST*	> ULN – 3*ULN	> 3*ULN – 5*ULN	> 5*ULN – 20*ULN	> 20*ULN
ALT *	> ULN – 3*ULN	> 3*ULN – 5*ULN	> 5*ULN – 20*ULN	> 20*ULN
GGT	> ULN – 2.5*ULN	>2.5*ULN – 5*ULN	> 5*ULN – 20*ULN	> 20*ULN
Bilirubin	> ULN – 1.5*ULN	>1.5*ULN – 3*ULN	> 3*ULN – 10*ULN	> 10*ULN
Fibrinogen^	%change of BL <25% or 0.75*LLN - < LLN	25%- <50% of BL or < 75*LLN – 0.5*LLN	50% - <75% of BL or < 0.5* LLN – 0.25*LLN	>= 75% of BL or < 50mg/dL or < 0.25*LLN
Calcium [mmol/L]*	2.0 - < LLN	1.75 - < 2.0	1.5 - < 1.75	< 1.5
Potassium [mmol/L]*	not defined	3.0 - < LLN	2.5 - < 3.0	< 2.5
Lipase	> ULN – 1.5*ULN	> 1.5*ULN – 2.0*ULN	> 2.0*ULN – 5.0*ULN	> 5.0*ULN
Amylase	> ULN – 1.5*ULN	> 1.5*ULN – 2.0*ULN	> 2.0*ULN – 5.0*ULN	> 5.0*ULN

BL: baseline value, LLN: Lower limit of normal, ULN: Upper limit of normal

^{^:} In case of conflicting criteria the higher grade will be assigned, % change only used when baseline is <LLN



^{*:} Clinical criteria from CTC AE 4.0 grading were not considered in order to assign grades